

```

Tyr Asp Lys Ser Ile Val Asp Ser Gly Thr Thr Asn Leu Arg Leu Pro
-      290              295              300
-
Lys Lys Val Phe Glu Ala Ala Val Lys Ser Ile Lys Ala Ala Ser Ser
-      305              310              315              320
-
Thr Glu Lys Phe Pro Asp Gly Phe Trp Leu Gly Glu Gln Leu Val Cys
-      325              330              335
-
Trp Gln Ala Gly Thr Thr Pro Trp Asn Ile Phe Pro Val Ile Ser Leu
-      340              345              350
-
Tyr Leu Met Gly Glu Val Thr Asn Gln Ser Phe Arg Ile Thr Ile Leu
-      355              360              365
-
Pro Gln Gln Tyr Leu Arg Pro Val Glu Asp Val Ala Thr Ser Gln Asp
-      370              375              380
-
Asp Cys Tyr Lys Phe Ala Ile Ser Gln Ser Ser Thr Gly Thr Val Met
-      385              390              395              400
-
Gly Ala Val Ile Met Glu Gly Phe Tyr Val Val Phe Asp Arg Ala Arg
-      405              410              415
-
Lys Arg Ile Gly Phe Ala Val Ser Ala Cys His Val His Asp Glu Phe
-      420              425              430
-
Arg Thr Ala Ala Val Glu Gly Pro Phe Val Thr Leu Asp Met Glu Asp
-      435              440              445
-
Cys Gly Tyr Asn Ile Pro Gln Thr Asp Glu Ser

```

450

455

<210> 25

<211> 1302

<212> DNA

<213> Homo sapiens

<400> 25

atgactcagc atggtattcg tctgccactg cgtagcggtc tgggtggcgc tccactgggt 60
 ctgcgtctgc cccgggagac cgacgaagag cccgaggagc cccggccggag gggcagcttt 120
 gtggagatgg tggacaacct gaggggcaag tcggggcagg gctactacg ggagatgacc 180
 gtgggcagcc ccccgagac gctcaacatc ctggtggata caggcagcag taactttgca 240
 gtgggtgtg cccccaccc cttcctgcat cgctactacc agaggcagct gtccagcaca 300
 taccgggacc tccggaagg tggttatgtg cctacaccc agggcaagt ggaaggggag 360
 ctgggcaccg acccggttaag catccccat ggcaccaacg tcaactgtgc tgccaacatt 420
 gctgccatca ctgaatcaga caagtcttc atcaacggct ccaactggga aggcattctg 480
 gggctggcct atgctgagat tgcaggcct gacgactccc tggagccttt ctttgnctct 540
 ctggttaagc agaccacgt tcccaacctc ttctccctgc acctttgtgg tgcctggctc 600
 cccctcaacc actctcaact gctgacctct gtcggaggga gcatgatcat tggaggtatc 660
 gaccactcgc tgtacacagg cagctctcgg tatacaccca tccggcggga gtggtattat 720
 gaggtcatca ttgtgcgggt ggagatcaat ggacaggatc tgaatatgga ctgcaaggag 780
 tacaactatg acaagagcat tgtggacagt ggcaccacra accttcgttt gcccaagaa 840
 gtgtttgaag ctgcagtcaa atccatcaag gcagcctcct ccacggagaa gttccctgat 900
 ggtttctggc taggagagca gctgggtgtc tggcaagcag gcaccacccc ttggaacatt 960
 ttcacgtca tctactcta cctaattgggt gaggttacca accagtcctt ccgcatcacc 1020
 atcttcgc agcaatacct ggggcagtg gaagatgtg ccacgtccca agacgactgt 1080
 tacaagttt ccattctaca gtcatccacg ggcactgtta tgggagctgt tatcatggag 1140
 ggcttctacg ttgtcttga cggggccga aaacgaattg gctttgctgt cagcgcttgc 1200
 catgtgcacg atgagttcag gacggcagcg gtggaaggcc cttttgtcac cttggacatg 1260
 gaagactgtg gctacaacat tccacagaca gatgagtcac ga 1302

```

-
<210> 26
-
<211> 433
-
<212> PRT
-
<213> Homo sapiens
-
-
<400> 26
Met Thr Gln His Gly Ile Arg Leu Pro Leu Arg Ser Gly Leu Gly Gly
1           5           10           15
-
Ala Pro Leu Gly Leu Arg Leu Pro Arg Glu Thr Asp Glu Glu Pro Glu
           20           25           30
-
Glu Pro Gly Arg Arg Gly Ser Phe Val Glu Met Val Asp Asn Leu Arg
           35           40           45
-
Gly Lys Ser Gly Gln Gly Tyr Tyr Val Glu Met Thr Val Gly Ser Pro
           50           55           60
-
Pro Gln Thr Leu Asn Ile Leu Val Asp Thr Gly Ser Ser Asn Phe Ala
           65           70           75           80
-
Val Gly Ala Ala Pro His Pro Phe Leu His Arg Tyr Tyr Gln Arg Gln
           85           90           95
-
Leu Ser Ser Thr Tyr Arg Asp Leu Arg Lys Gly Val Tyr Val Pro Tyr
           100          105          110
-
Thr Gln Gly Lys Trp Glu Gly Glu Leu Gly Thr Asp Leu Val Ser Ile
           115          120          125
-
-

```